

REPLACEMENT SHEET

FIG. 13

ROW	PRIMER SEQUENCE	SEQ ID NO	SEQUENCED SEQUENCE	SEQ ID NO	PREDICTED GAM RNA	SEQ ID NO	DIST-ANCE	GAM NAME
1*	AATTGCTTGAAC	4254671	CCAGGAAGTGGG	4254700	AATTGCTTGAACCCAGGAAGTGGG	4254729	0	25-A
2*	ACTGCCTCC	4254672	AGCCTGGGC	4254701	ACTGCCTCCAGCCTGGGCTAC	4254730	0	351661-A
3	CACTGCACTC	4254673	CAGCCGAGCAACA	4254702	CATGCACCTCCAGCCGAGCAA	4254731	0	351946-A
4	CTAGACTGAAG	4254674	CTCCTTGAGGAC	4254703	CTAGACTGAAGCTCTTGAGGA	4254732	0	352759-A
5	GAAGTTGAAG	4254675	CCTGTTGTICA	4254704	GAAGTTGAAGCCTGTGTTCA	4254733	0	4426-A
6	TCACITGCAAC	4254676	CTCCACCA	4254705	(TCACTGCAACCTCCACCACGTG), (TCACTGCAACCTCCACCACGCT)	4254734, 4254735	0	(357950-A), (352721-A)
7*	TCTAGAGAAAG	4254677	GAAGTTCAGA	4254706	TCTAGAGAAAGGAAGTTCAGA	4254736	0	337950-A
8	GGGCAGTGGG	4254678	GCTGGAA	4254707	GGGCGTGGAGCTGGATGATGT	4254737	1	351996-A
9	AATTGCTTGAAC	4254679	CCAAGAAGTGGG	4254708	AATCACTTGAACCCCAAGAAGTG	4254738	2	351874-A
10	AGCAGGCCA	4254680	GGGTTTGT	4254709	AGCAAGACCAGGGTTTGTGT	4254739	2	352083-A
11	AGGCAAGACG	4254681	GACCAGA	4254710	AGGCAGAGAGACCAGAGACT	4254740	2	351944-A
12	AGGGAAGAAT	4254682	TAATGTGAA	4254711	GGGAATAATTAATGTGAAGTC	4254741	2	353325-A
13	AGGGAAGAAT	4254683	TAATGTGAG	4254712	AGGGAATAATTAATGTGAAGTC	4254742	2	352649-A
14	ATTCAGTTG	4254684	CCCAIGTTT	4254713	(ATTCAATGCCCAATGTTT), (TATTCATGCCCAATGTTGTA)	4254743, 4254744	2	(352957-A), 352960-A
15	CTAGACTGAAG	4254685	CTCCTTGAGG	4254714	CTGGACTGAGCTCCTTGAGGCC	4254745	2	352288-A
16	TTCAGAGTGGT	4254686	TAAGTTCTG	4254715	TTCAGAGTGGTAAAGTTCTGTCA	4254746	2	353875-A
17	TTCAGAGTGGT	4254687	TAAGTTCTGC	4254716	TTCAGAGTGGTAAAGTTCTGTCTT	4254747	2	351940-A
18	AGCAGGCCA	4254688	GAAGGAAG	4254717	AGGCCAAGGAAGGAAGCAGAGG	4254748	3	352496-A
19	AGTTTGCCITG	4254689	TAAGAAAG	4254718	AGTTTGTTAAGAAAGC	4254749	3	352518-A
20	ATCAGAGGGTG	4254690	GGTGCTAA	4254719	ATTAGGAGAGTGGGTGCTAAGT	4254750	3	352511-A
21	ATGGTGGAG	4254691	AGTTTCTCAGT	4254720	TGGAGGAGAGTTTGTCAAGTATAG	4254751	3	353484-A
22	CCAGGAAG	4254692	TGGAGCCTGGGC	4254721	CCCGGTGGAGCCTGGGCTGTG	4254752	3	351990-A
23	GGCAGTGGG	4254693	GGTCCGT	4254722	AGGCAGGAGGTCGTCCTCTTC	4254753	3	353880-A
24	GGCAGTGGG	4254694	TCTAGAC	4254723	GTGACAGTGAATCTAGACAGAC	4254754	3	352810-A
25	TCAAGCTCATT	4254695	CACTAAA	4254724	CTCAGCTCATCCCAATAATCCC	4254755	3	353184-A
26	TGGAAGTT	4254696	GGTTGTATGGTT	4254725	GGATGGTGGTGTATGGTTG	4254756	3	353855-A
27	TGGAGAGTT	4254697	CCATATTTG	4254726	TGATAGATCCATATTTGGTAA	4254757	3	352004-A
28	TGGAGAGTT	4254698	GTTGTACAGT	4254727	TGGGTTTGTGTTGACAGTGA	4254758	3	353160-A
29	TCACITGCAAC	4254699	CTCCACC	4254728	TCACTGCAACCTCCACCCTTCCG	4254759	0	353856-A